

iGEM 2016: Pittsburgh  
**Week 5 Lab Notebook**

*Monday, June 20*

Lab meeting

Double digest of T7-RBS -- lacZ/eGFP to check for insert (Praneeth)

0.5 uL plasmid

0.5 uL EcoRI

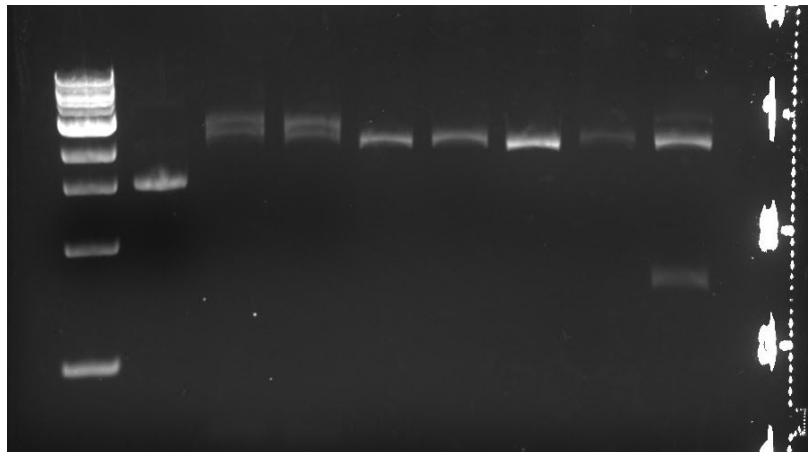
0.5 uL SpeI

1 uL 10x buffer

7.5 uL H<sub>2</sub>O

Gel electrophoresis check

Lane	1	ladder
	2	control
	3	T7 LacZ 1
	4	T7 LacZ 2
	5	T7 eGFP 1
	6	T7 eGFP 2
	7	T7 eGFP 3
	8	T7 eGFP 4
	9	T7 eGFP 5



Transformation of Collins plasmids (Claire)

D\_LacZ (13), kan resistance

pAG\_TS2AT05 (14), amp resistance

G\_LacZ (15), kan resistance

pAG\_TS2AT03 (16), amp resistance

Sequence of T7-RBS -- amilCP not convincing for amilCP insertion direction

Correct sequence from wrong primer--misabeled primers?

Double digest of 3-2 and 3-5 T7-RBS -- amilCP with EcoRI and SpeI

Sign up for InterLab study

Email Ronald Joseph (Pittsburgh Public Schools water testing) [rj1@pghboe.net](mailto:rj1@pghboe.net)

Tuesday, June 21

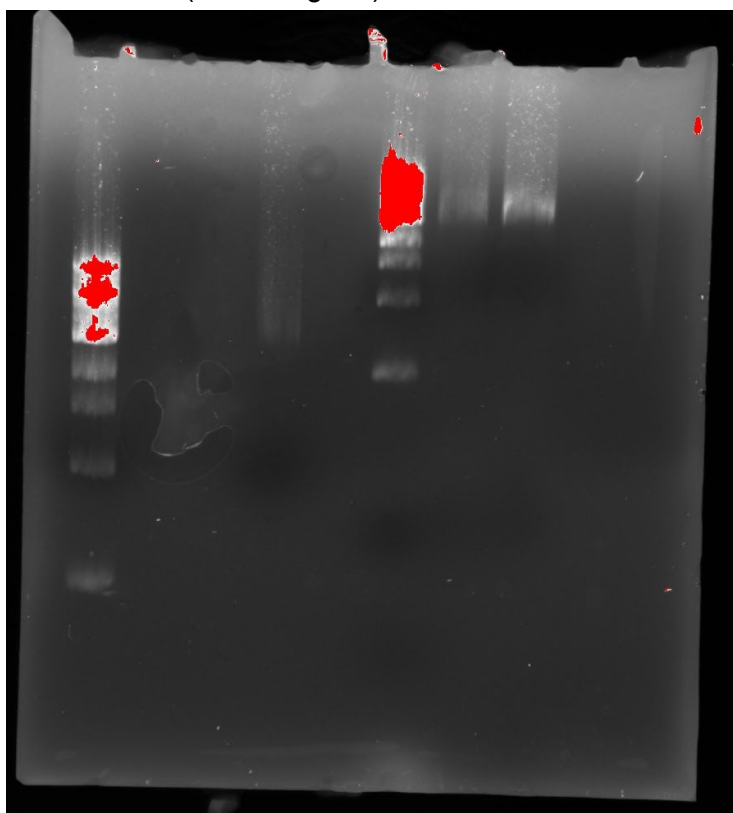
[Gel electrophoresis](#) of double digest 3-2 and 3-5 T7-RBS -- amilCP (Claire)

1% agarose gel

Lane	1	ladder
	2	3-2 (1 uL)
	3	3-5 (1 uL)
	4	uncut 3-5 control

Couldn't see lanes 2 and 3, run again

Lane	6	ladder
	7	3-2 (remaining rxn)
	8	3-5 (remaining rxn)



Did not confirm amilCP direction

[Double digest](#) of 3-2 T7-RBS -- amilCP (Maddie)

~1 uL SpeI (left over)

1 uL EcoRI

2 ug DNA = 15.34 uL DNA

2 uL 10x buffer

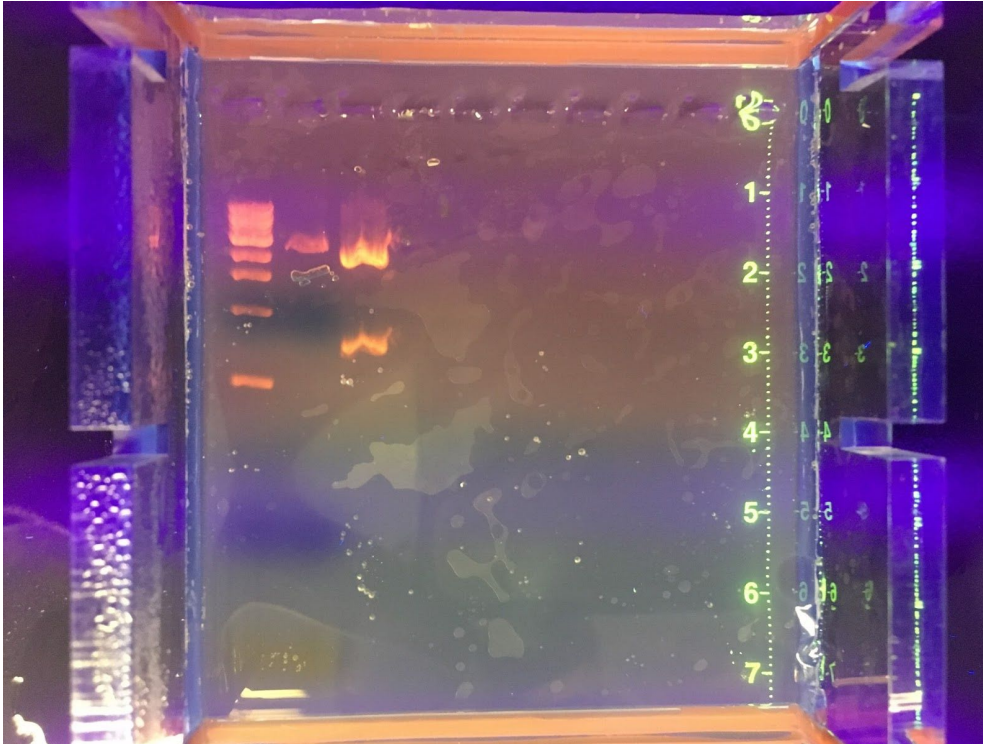
.66 uL H<sub>2</sub>O

No heat inactivation

[Gel electrophoresis](#) of digested 3-2 T7-RBS -- amilCP (Maddie)

Lane	1	ladder
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- 2 undigested plasmid
- 3 digest reaction



[Liquid cultures](#) of Collins plasmids (2 of each plasmid), into shaker at 2 pm (Claire)



Wednesday, June 22

[Ligation](#) of T7-RBS -- amilCP to terminator (Claire, Maddie)

Control	1 uL	terminator
	0 uL	T7-RBS -- amilCP
	2 uL	buffer
	16 uL	H <sub>2</sub> O
	1 uL	T4 ligase
3:1	1 uL	terminator
	50 ng= 4.76 uL	T7-RBS -- amilCP
	2 uL	buffer

11.24 uL H<sub>2</sub>O  
 1 uL T4 ligase  
 x:1 1 uL terminator  
 5.76 uL T7-RBS -- amilCP  
 2 uL buffer  
 10.76 uL H<sub>2</sub>O  
 1 uL T4 ligase

Transformation of ligation reaction (Claire)

5 uL of each reaction into 66.6 uL of Cheryl's competent cells  
 Ice 10 min  
 42 degC 2 min  
 Ice 5 min  
 200 uL SOC media  
 Shaker at 37 degC for 1 hr. 45 min.  
 Into incubator at 12:45 pm

Mini-prep Collins plasmids liquid cultures (Maddie, Praneeth)

Concentrations (ng/uL)

	<b>13</b>	<b>14</b>	<b>15</b>	<b>16</b>
<b>1</b>	81.3	162.5	95.4	141.9
<b>2</b>	83.6	129.1	71.6	148.6

Cell-free extract volume optimization (Maddie, Praneeth)

T7--GFP (concentration = 120.8 ng/uL)

DNA amount = 5 ng/uL

	<b>25 µL</b>	<b>25 Control</b>	<b>20 µL</b>	<b>20 C</b>	<b>10 µL</b>	<b>10 C</b>	<b>5 µL</b>	<b>5 C</b>
<b>Solution A</b>	10	10	8	8	4 µL	4 µL	2 µL	2 µL
<b>Solution B</b>	7.5	7.5	6	6	3 µL	3 µL	1.5 µL	1.5 µL
<b>RNAse</b>	0.125	0.125	0.2	0.2	0.1 µL	0.1 µL	0.05	0.05
<b>H<sub>2</sub>O</b>	6.345	7.355	4.972	5.8	2.486	2.9	1.243	1.45
<b>DNA</b>	1.03	0	0.828	0	0.414	0	0.207	0

*New RNAse at twice the concentration of old RNAse*

\* all 25 uL and 20 uL reactions, including controls, were made from Solutions A and B that were mixed on June 17, then frozen at -80 degC because RNAse was low

Incubate 2 hours at 37 degC

Top off all reactions with water to 25 uL (min volume of plate reader)

	1	2	3	4	5	6	7	8	9	10	11	12
A					5 C	5-1	5-2	10 C	10-1	10-2	20 C	20-1
B	20-2	25 C	25-1	25-2								

Results (rfu) 485 emission, 585 excitation

Read multiple times because wasn't sure if plate reader settings were correct

	1	2	3	4	5	6	7	8	9	10	11	12
A	3.721	5.346	3.688	3.716	3.274	34.258	47.955	4.155	31.730	5.128	4.421	5.827
B	31.660	6.907	3.261	3.594	3.842	4.136	3.658	3.850	3.478	3.973	4.316	7.949

	1	2	3	4	5	6	7	8	9	10	11	12
A	0.330	0.946	0.299	0.448	0.361	26.984	37.667	0.459	0.749	0.878	0.689	1.093
B	21.106	0.728	0.729	0.720	0.321	0.328	0.355	0.260	0.279	0.298	0.487	1.041

	1	2	3	4	5	6	7	8	9	10	11	12
A	0.327	0.953	0.340	0.389	0.426	27.122	37.081	0.536	0.846	1.011	0.559	0.861
B	21.625	0.916	1.008	0.902	0.264	0.308	0.346	0.266	0.243	0.276	0.320	1.079

Only 5-mL worked. Forgot to add DNA? Refrozen cell extract?

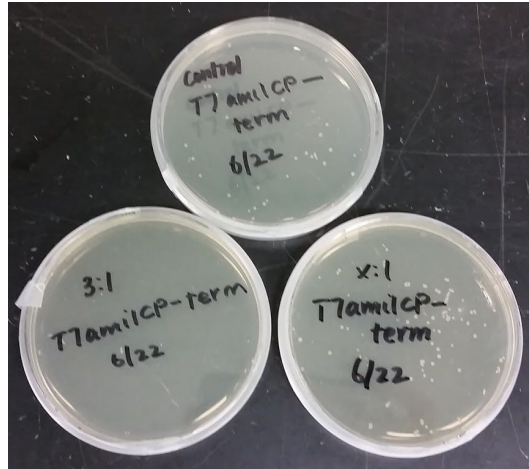
Linear T7-RBS -- amilCP in [cell extract](#) (Aife)

T7-RBS -- amilCP digested with EcoRI, concentration = 52.16 ng/uL

	5 uL	5 uL control	1 uL	1 uL control
<b>Sol'n A</b>	2	2	0.4	0.4
<b>Sol'n B</b>	1.5	1.5	0.3	0.3
<b>RNAse</b>	0.25	0.25	0.05	0.05
<b>DNA</b>	0.96	0	0.19	0
<b>water</b>	0.29	1.25	0.06	0.25

Thursday, June 23

Colonies grew on x:1 and control plates (but more on x:1) for T7amilCP--term



Liquid cultures of 5 colonies from x:1 T7amilCP--term plate (Praneeth)

Digest T7-eGFP 2, 3, 4, 5 with EcoRI and SpeI (Claire)

1 uL SpeI

1 uL EcoRI

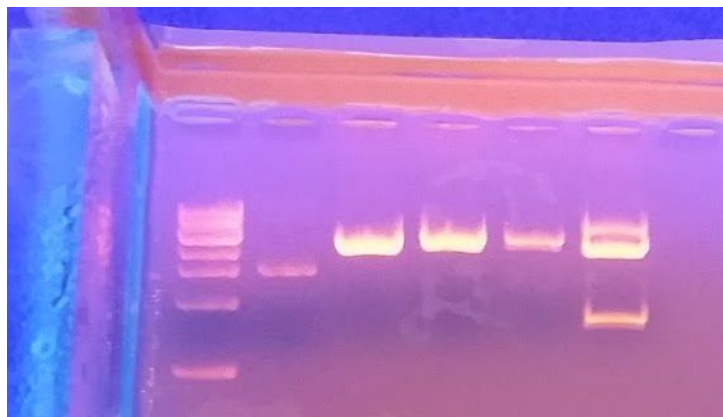
2 ug DNA = 13.24 uL T7-eGFP 2  
10.07 uL T7-eGFP 3  
10.94 uL T7-eGFP 4  
9.58 uL T7-eGFP 5

2 uL buffer

Water = 2.76 uL 2  
5.93 uL 3  
5.06 uL 4  
6.42 uL 5

Gel electrophoresis T7-eGFP (Claire)

Lane	1	ladder
	2	uncut T7-eGFP 2
	3	T7-eGFP 2
	4	T7-eGFP 3
	5	T7-eGFP 4
	6	T7-eGFP 5 (extract smallest band)



Gel extraction of lane 6 (T7-eGFP 5) (Claire)

Concentration = 23.1 ng/uL

Ligation T7-eGFP -- terminator (Claire)

Control      1 uL    terminator (double digested)  
                 0 uL    T7-eGFP  
                 2 uL    buffer  
                 16 uL   H2O  
                 1 uL    ligase  
3:1            1 uL    terminator (double digested)  
                 50 ng = 2.16 uL T7-eGFP  
                 2 uL    buffer  
                 13.84 uL H2O  
                 1 uL    ligase  
7:1            1 uL    terminator (double digested)  
                 115 ng = 4.98 uL T7-eGFP  
                 2 uL    buffer  
                 11.02 uL H2O  
                 1 uL    ligase

Transformation: 5 uL of each reaction into 66.6 uL Cheryl's cells (Claire)

Collins plasmids in cell-free extract (Praneeth)

Substrate: Chlorophenol Red- $\beta$ -D-galactopyranoside at 12 mg/mL (20x)  
working concentration = 0.6 mg/mL

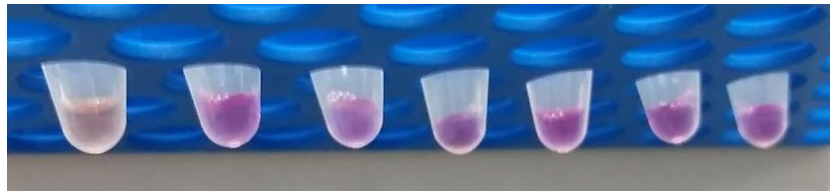
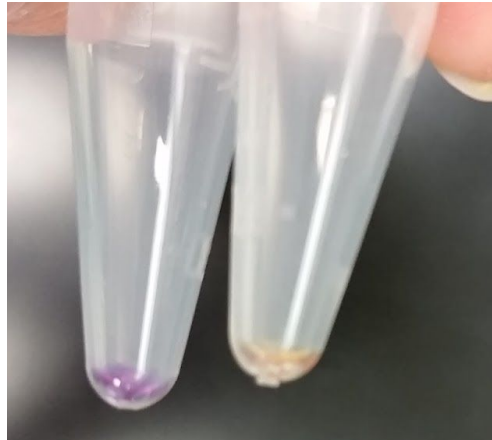
Switch: D\_LacZ (13)    Trigger: pAG\_Ts2AT05 (14)

Amt trigger	50 ng	25 ng	10 ng	0 ng
Solution A	2	2	2	2
Solution B	1.5	1.5	1.5	1.5
RNAse	0.25	0.25	0.25	0.25
Switch (13-2) 50 ng	0.60	0.60	0.60	0.60
Trigger (14-1)	0.31	0.15	0.06	0.00
Substrate	0.25	0.25	0.25	0.25
Water	0.09	0.25	0.34	0.40

Switch: G\_LacZ (15) Trigger: pAG\_Ts2AT03 (15)

Amt trigger	50 ng	25 ng	10 ng	0 ng
<b>Solution A</b>	2	2	2	2
<b>Solution B</b>	1.5	1.5	1.5	1.5
<b>RNAse</b>	0.25	0.25	0.25	0.25
<b>Switch (15-1) 50 ng</b>	0.52	0.52	0.52	0.52
<b>Trigger (16-2)</b>	0.34	0.17	0.07	0.00
<b>Substrate</b>	0.25	0.25	0.25	0.25
<b>Water</b>	0.14	0.31	0.41	0.48

Incubate 2 hours at 37 degC; color change visible around 1 hour



D plasmids (13 and 15), increasing concentration from left to right



G plasmids (14 and 16), increasing concentration from left to right

Clear plate:

	1	2	3	4	5	6	7
A	13-0	13-10 1	13-10 2	13-25 1	13-25 2	13-50 1	13-50 2
B	15-0	15-10 1	15-10 2	15-25 1	15-25 2	15-50 1	15-50 2

Add 20 uL water to each well to achieve plate reader's min concentration

Wavelength = 570 nm

	1	2	3	4	5	6	7
A	0.122	0.449	0.250	0.373	0.378	0.437	0.406
B	1.048	0.420	0.592	0.573	0.434	0.587	0.555

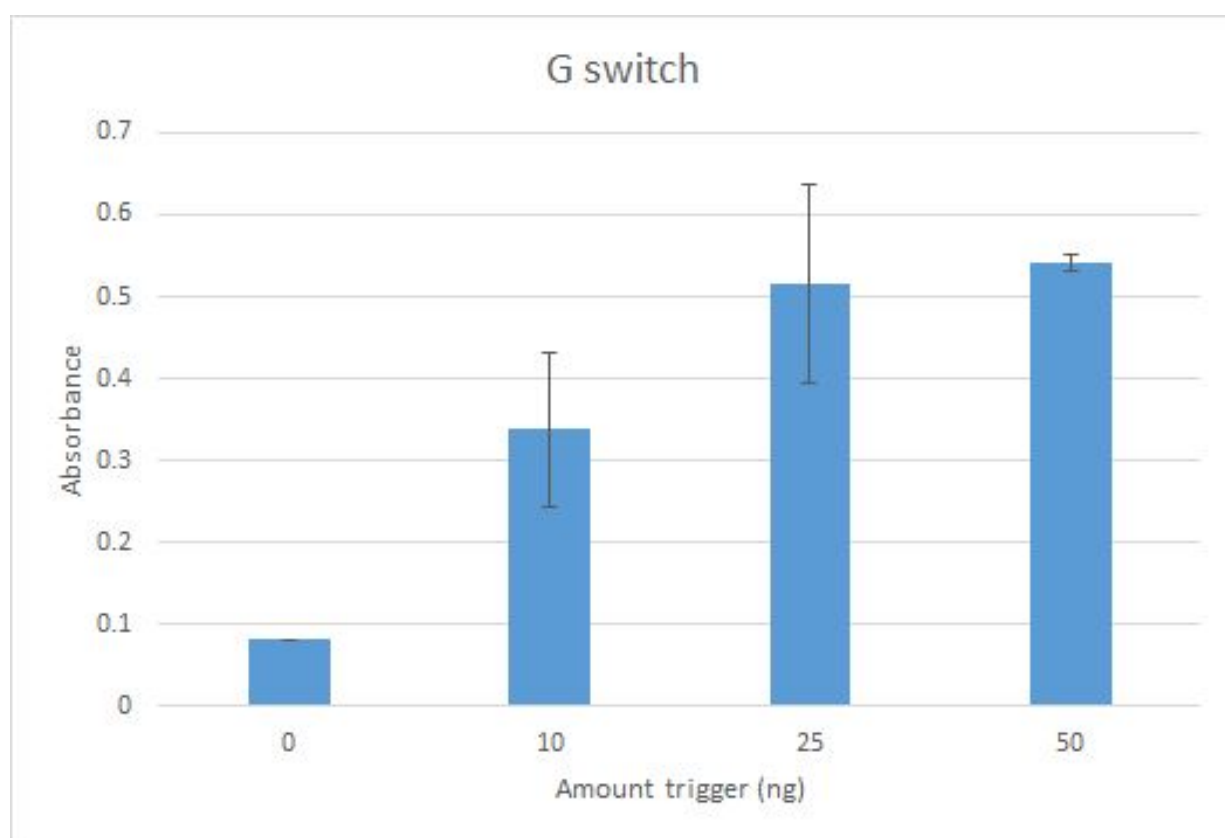
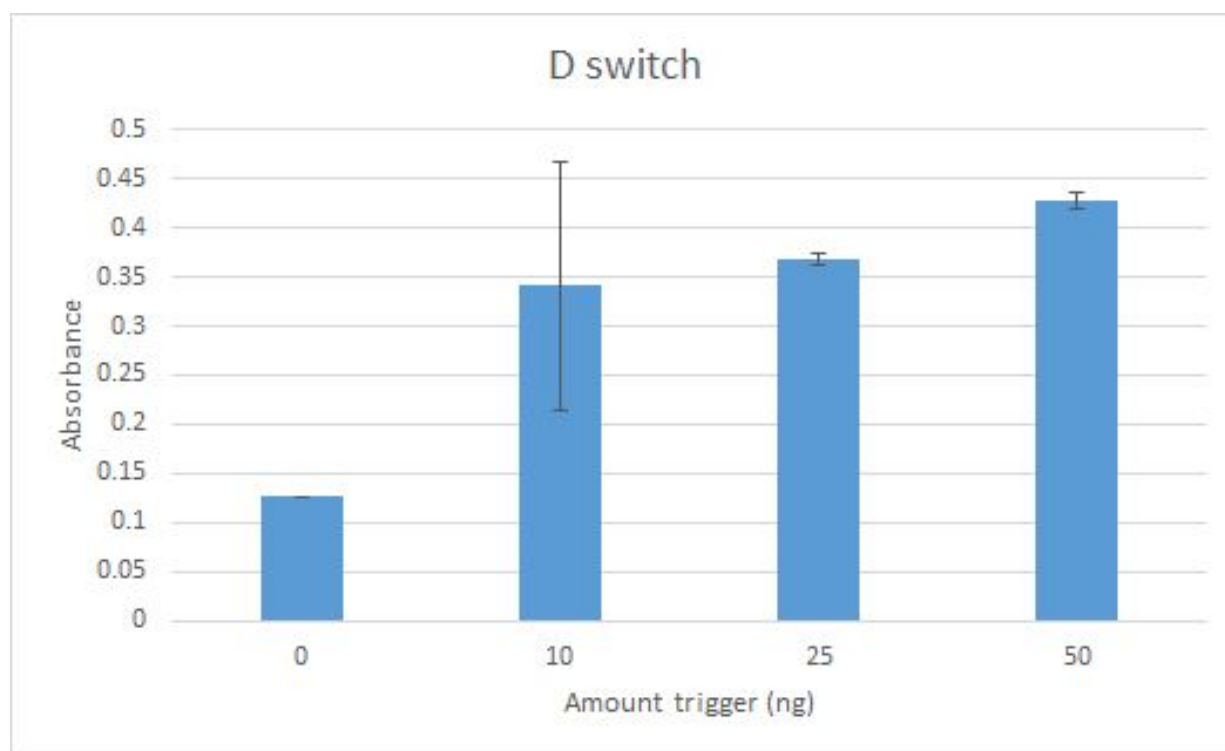
Wells not evenly covered, so add 25 uL more water (45 uL total)

Results from Roy lab plate reader similar, so Roy lab reader good for LacZ

	1	2	3	4	5	6	7
A	0.127	0.440	0.255	0.369	0.379	0.435	0.422
B	0.565	0.409	0.578	0.316	0.430	0.543	0.551

Pipette out bubbles from B1 and B3, read at Roy lab plate reader

	1	2	3	4	5	6	7
A	0.126	0.431	0.252	0.364	0.372	0.434	0.421
B	0.082	0.404	0.272	0.600	0.429	0.534	0.548



Error bars represent standard deviation

Inequivalence of samples of same concentration may be due to pipetting error--very small volumes of DNA (e.g., 0.06 uL) are difficult to achieve precision with. Next time, dilute DNA.



Friday, June 24

T7-eGFP -- terminator plates (growth on 3:1, none on 7:1, none on control!)



Mini-prep T7amilCP--terminator liquid cultures

Concentrations (ng/uL):	1	109.7
	2	167.9
	3	130.6
	4	175.5
	5	170.3

Digest mini-prepped DNA to check for ligations with EcoRI and PstI

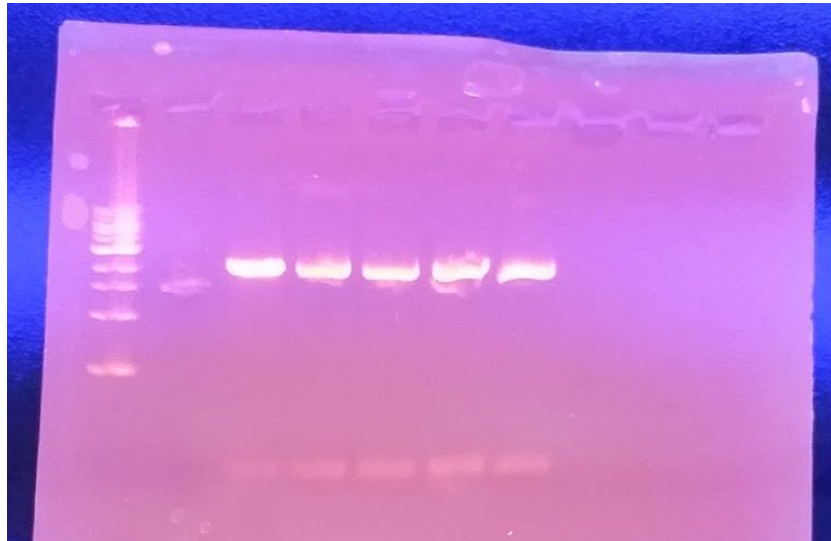
0.5 uL EcoRI

0.5 uL PstI		
1 ug DNA =	9.12 uL	1
	5.96 uL	2
	7.66 uL	3
	5.70 uL	4
	5.87 uL	5
1 uL buffer		
H2O =	0 uL	1
	2.04 uL	2
	0.34 uL	3
	2.30 uL	4
	2.13 uL	5

Gel electrophoresis of digests

Lane	1	2 uL ladder + 0.5 uL loading dye
	2	2 uL MLK terminator (uncut) + 0.5 uL loading dye
	3	1
	4	2
	5	3
	6	4
	7	5

No successful ligations?



LB + CM plates

Sunday, June 26

Liquid cultures (Maddie)

x:1 T7 amilCP -- terminator (x5)

3:1 T7 eGFP -- terminator (x5)